HARBERD et al Appln. No.: 10/809,945

August 9, 2004

Figure 12b shows a partial amino acid sequence (SEQ ID NO:11) of the wheat rht-10 allele.

Please replace the paragraph beginning at page 45, 7000 line 3, with the following rewritten paragraph:

Previously, we cloned the GAI gene of Arabidopsis (PCT/GB97/00390 - WO97/29123 published 14 August 1997). Comparison of the DNA sequences of the wild-type (GAI) and mutant (gai) alleles showed that gai encodes a mutant predicted protein product (qai) which lacks a segment of 17 amino acids from close to the N-terminus of the protein. Screening of the DNA sequence databases with the GAI sequence revealed the existence of a rice EST (D39460) which contains a region of sequence very closely related to that of the segment that is deleted from GAI in the gai protein. A comparison of the predicted amino acid sequences from the region DELLA (SEQ ID NO:107) to EQLE (SEQ ID NO:108) are identical in both sequences. The two differences (V/A; E/D) are conservative substitutions, in which one amino acid residue is replaced by another having very similar chemical properties. In addition, the region of identity extends beyond the boundary of the deletion